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RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer-readable form:

Application Serial Number: 09/634,252

Source: 1652

Date Processed by STIC: 12-20-00

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DEC 27 2000

ENTER 1842200

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-306-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT MARK SPENCER, 703-306-4216.

PATENTIN 2.1 e-mail help: patin21help@uspto.gov or phone 703-306-4119 (R. Wax)

PATENTIN 3.0 e-mail help: patin30help@uspto.gov or phone 703-306-4119 (R. Wax)

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER
VERSION 3.0 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND
TRADEMARK OFFICE WEBSITE. SEE BELOW.

Checker Version 3.0

The Checker Version 3.0 application is a state-of-the-art Windows based software program employing a logical and intuitive user-interface to check whether a sequence listing is in compliance with format and content rules. Checker Version 3.0 works for sequence listings generated for the original version of 37 CFR §§1.821 - 1.825 effective October 1, 1990 (old rules) and the revised version (new rules) effective July 1, 1994 as well as World Intellectual Property Organization (WIPO) Standard ST.25.

Checker Version 3.0 replaces the previous DOS-based version of Checker, and is V2K compliant. Checker allows public users to check sequence listings in Computer Readable form.

Use of Checker prior to filing the sequence listing is recommended.

Checker Version 3.0 can be downloaded from the U.S. Patent and Trademark Office website at:
<http://www.uspto.gov/web/offices/tac/checker>

1652

RAW SEQUENCE LISTING DATE: 12/20/2000
 PATENT APPLICATION: US/09/634,252 TIME: 10:53:44

Input Set : A:\SVPH3137.txt.txt
 Output Set: N:\CRF3\12202000\I634252.raw

3 <110> APPLICANT: Cerretti, Douglas P.
 5 <120> TITLE OF INVENTION: SVPH3-13 AND SVPH3-17 DNA AND POLYPEPTIDES
 7 <130> FILE REFERENCE: 03260.0051-00304
 C--> 9 <140> CURRENT APPLICATION NUMBER: US/09/634,252
 C--> 10 <141> CURRENT FILING DATE: 2000-08-07
 12 <150> PRIOR APPLICATION NUMBER: 60/074,310
 13 <151> PRIOR FILING DATE: 1998-02-11
 15 <160> NUMBER OF SEQ ID NOS: 10
 17 <170> SOFTWARE: PatentIn Ver. 2.0

Does Not Comply
 Corrected Diskette Needed

ERRORED SEQUENCES

478 <210> SEQ ID NO: 7
 479 <211> LENGTH: 34
 480 <212> TYPE: DNA
 481 <213> ORGANISM: Homo sapiens
 483 <400> SEQUENCE: 7
 E--> 484 tatcttccta aaaagttcac ccagtgtaa attg
 486 <210> SEQ ID NO: 8
 487 <211> LENGTH: 27
 488 <212> TYPE: DNA
 489 <213> ORGANISM: Homo sapiens
 491 <400> SEQUENCE: 8
 E--> 492 tgctttctat tcccatcga atagccc

Number of bases differ:
 34 listed
 33 shown

34

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27

27 listed
 26 shown

VERIFICATION SUMMARY

PATENT APPLICATION: US/09/634,252

DATE: 12/20/2000

TIME: 10:53:45

Input Set : A:\SVPH3137.txt.txt

Output Set: N:\CRF3\12202000\I634252.raw

L:9 M:270 C: Current Application Number differs, Replaced Application Number
L:10 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:484 M:254 E: No. of Bases conflict, LENGTH:Input:34 Counted:33 SEQ:7
L:484 M:252 E: No. of Seq. differs, <211>LENGTH:Input:34 Found:33 SEQ:7
L:492 M:254 E: No. of Bases conflict, LENGTH:Input:27 Counted:26 SEQ:8
L:492 M:252 E: No. of Seq. differs, <211>LENGTH:Input:27 Found:26 SEQ:8